



Replacement Sheet

A.

1	human, ZAKI-4 (hCsp2)	1	EFVDPVRPRVRLGQQA S I P E D D O O L F L C C I D R D W A V T Q C F A E E A F Q A L T D F S - D L P N S T F A C R V H Q S V F E E E S E S K E P K E	1	---DYS-TVACVVDVVFTHNQVVRKFPK	1	PKPK
1	murine, ZAKI-4 (mCsp2)	1	EFVDPVRPRVRLGQQA S I P E D D O O L F L C C I D R D W A V T Q C F A E E A F Q A L T D F S - D L P N S T F A C R V H Q S V F E E E S E S K E P K E	1	---DYS-TVACVVDVVFTHNQVVRKFPK	1	PKPK
1	human, D9CRI (hCsp1)	1	EFVDPVRPRVRLGQQA S I P E D D O O L F L C C I D R D W A V T Q C F A E E A F Q A L T D F S - D L P N S T F A C R V H Q S V F E E E S E S K E P K E	1	---DYS-TVACVVDVVFTHNQVVRKFPK	1	PKPK
1	hamster, Adapt78	1	EFVDPVRPRVRLGQQA S I P E D D O O L F L C C I D R D W A V T Q C F A E E A F Q A L T D F S - D L P N S T F A C R V H Q S V F E E E S E S K E P K E	1	---DYS-TVACVVDVVFTHNQVVRKFPK	1	PKPK
1	murine, D9CRI (mCsp1)	1	EFVDPVRPRVRLGQQA S I P E D D O O L F L C C I D R D W A V T Q C F A E E A F Q A L T D F S - D L P N S T F A C R V H Q S V F E E E S E S K E P K E	1	---DYS-TVACVVDVVFTHNQVVRKFPK	1	PKPK
1	C. elegans	1	EFVDPVRPRVRLGQQA S I P E D D O O L F L C C I D R D W A V T Q C F A E E A F Q A L T D F S - D L P N S T F A C R V H Q S V F E E E S E S K E P K E	1	---DYS-TVACVVDVVFTHNQVVRKFPK	1	PKPK
1	S. cerevisiae	1	EFVDPVRPRVRLGQQA S I P E D D O O L F L C C I D R D W A V T Q C F A E E A F Q A L T D F S - D L P N S T F A C R V H Q S V F E E E S E S K E P K E	1	---DYS-TVACVVDVVFTHNQVVRKFPK	1	PKPK
1	S. pombe	1	EFVDPVRPRVRLGQQA S I P E D D O O L F L C C I D R D W A V T Q C F A E E A F Q A L T D F S - D L P N S T F A C R V H Q S V F E E E S E S K E P K E	1	---DYS-TVACVVDVVFTHNQVVRKFPK	1	PKPK
29	human, ZAKI-4 (hCsp2)	29	GLERTYDDC V T F O L E K S F R K R V R I N F S N P K S A A R A R L E L H E T Q E R G K K ---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	29	---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	29	PKPK
80	murine, ZAKI-4 (mCsp2)	80	GLERTYDEC V T F O L E K S F R K R V R I N F S N P K S A A R A R L E L H E T Q E R G K K ---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	80	---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	80	PKPK
8	human, D9CRI (hCsp1)	8	GLERTYDDC V T F O L E K S F R K R V R I N F S N P K S A A R A R L E L H E T Q E R G K K ---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	8	---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	8	PKPK
34	hamster, Adapt78	34	GLERTYDDC V T F O L E K S F R K R V R I N F S N P K S A A R A R L E L H E T Q E R G K K ---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	34	---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	34	PKPK
53	murine, D9CRI (mCsp1)	53	GLERTYDDC V T F O L E K S F R K R V R I N F S N P K S A A R A R L E L H E T Q E R G K K ---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	53	---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	53	PKPK
33	C. elegans	33	GLERTYDDC V T F O L E K S F R K R V R I N F S N P K S A A R A R L E L H E T Q E R G K K ---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	33	---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	33	PKPK
38	S. cerevisiae	38	GLERTYDDC V T F O L E K S F R K R V R I N F S N P K S A A R A R L E L H E T Q E R G K K ---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	38	---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	38	PKPK
16	S. pombe	16	GLERTYDDC V T F O L E K S F R K R V R I N F S N P K S A A R A R L E L H E T Q E R G K K ---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	16	---K L Y F A Q V Q T P E N D Q D K L H L A P P O P A K Q F L	16	PKPK
106	human, ZAKI-4 (hCsp2)	106	ISPPSPSP ---V G W K P E D A T P V N Y D L L Y A A K L G P G E K Y E L H A T T P S V V V H V C S D E E E P K ---T S ---P K P K	106	---T S ---P K P K	106	PKPK
157	murine, ZAKI-4 (mCsp2)	157	ISPPSPSP ---V G W K P E D A T P V N Y D L L Y A A K L G P G E K Y E L H A T T P S V V V H V C S D E E E P K ---T S ---P K P K	157	---T S ---P K P K	157	PKPK
81	human, D9CRI (hCsp1)	81	ISPPASPP ---V G W K Q V E D A T P V N Y D L L Y A A K L G P G E K Y E L H A T T P S V V V H V C S D E E E P K ---T S ---P K P K	81	---T S ---P K P K	81	PKPK
107	hamster, Adapt78	107	ISPPASPP ---V G W K Q V E D A T P V N Y D L L Y A A K L G P G E K Y E L H A T T P S V V V H V C S D E E E P K ---T S ---P K P K	107	---T S ---P K P K	107	PKPK
107	murine, D9CRI (mCsp1)	107	ISPPASPP ---V G W K Q V E D A T P V N Y D L L Y A A K L G P G E K Y E L H A T T P S V V V H V C S D E E E P K ---T S ---P K P K	107	---T S ---P K P K	107	PKPK
128	C. elegans	128	ISPPASPP ---V G W K Q V E D A T P V N Y D L L Y A A K L G P G E K Y E L H A T T P S V V V H V C S D E E E P K ---T S ---P K P K	128	---T S ---P K P K	128	PKPK
112	S. cerevisiae	112	ISPPASPP ---V G W K Q V E D A T P V N Y D L L Y A A K L G P G E K Y E L H A T T P S V V V H V C S D E E E P K ---T S ---P K P K	112	---T S ---P K P K	112	PKPK
96	S. pombe	96	ISPPASPP ---V G W K Q V E D A T P V N Y D L L Y A A K L G P G E K Y E L H A T T P S V V V H V C S D E E E P K ---T S ---P K P K	96	---T S ---P K P K	96	PKPK
178	human, ZAKI-4 (hCsp2)	178	IIQTR ---R P ---G L P P G S M ---	178	---G L P P G S M ---	178	PKPK
227	murine, ZAKI-4 (mCsp2)	227	IIQTR ---R P ---G L P P G S M ---	227	---G L P P G S M ---	227	PKPK
157	human, D9CRI (hCsp1)	157	IIQTR ---R P ---G L P P G S M ---	157	---G L P P G S M ---	157	PKPK
183	hamster, Adapt78	183	IIQTR ---R P ---G L P P G S M ---	183	---G L P P G S M ---	183	PKPK
184	murine, D9CRI (mCsp1)	184	IIQTR ---R P ---G L P P G S M ---	184	---G L P P G S M ---	184	

Figure 1



B.

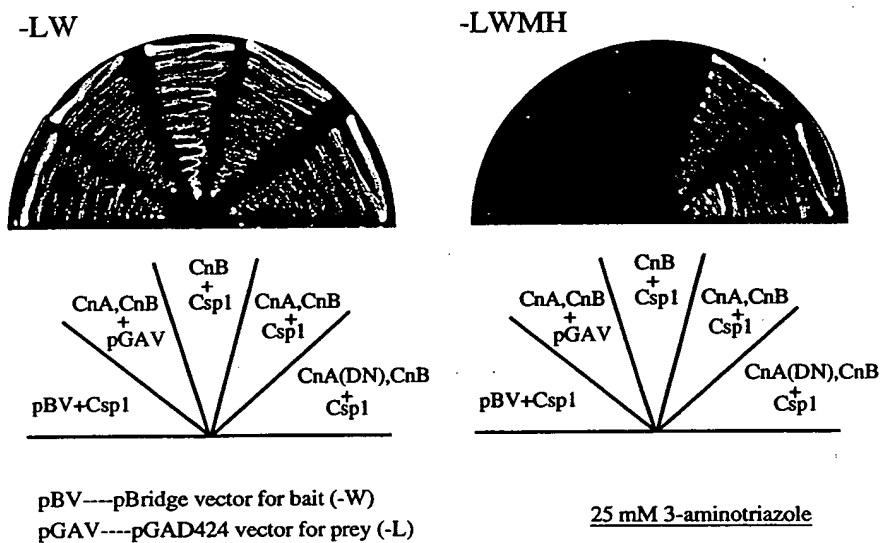


Figure 1 (continued)